

## SEQUENCE LISTING

<110> Kudoh, Masatake  
Yamamoto, Hiroaki

<120> (R)-2-OCTANOL DEHYDROGENASES, METHODS  
FOR PRODUCING THE ENZYMES, DNA ENCODING THE ENZYMES, AND  
METHODS FOR PRODUCING ALCOHOLS USING THE ENZYMES

<130> 06501-090001

<140> 09/978,758  
<141> 2001-10-16

<150> PCT/JP01/01082  
<151> 2001-02-15

<150> JP 2000-374593  
<151> 2000-12-08

<150> JP 2000-43506  
<151> 2000-02-16

<160> 20

<170> FastSEQ for Windows Version 4.0

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tca gga atc ggc tta agc gtc gca aaa aag ttc ctt cag ctc ggc gcc 96  
Ser Gly Ile Gly Leu Ser Val Ala Lys Lys Phe Leu Gln Leu Gly Ala  
20 25 30

aaa gta acg atc tct gat gtc agt gga gag aaa aaa tat cac gag act 144  
Lys Val Thr Ile Ser Asp Val Ser Gly Glu Lys Lys Tyr His Glu Thr  
35 40 45

gtt gtt gct ctg aaa gcc caa aat ctc aac act gac aac ctc cat tat 192  
Val Val Ala Leu Lys Ala Gln Asn Leu Asn Thr Asp Asn Leu His Tyr  
50 55 60

gta cag gca gat tcc agc aaa gaa gaa gat aac aag aaa ttg att tcg 240

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Val Gln Ala Asp Ser Ser Lys Glu Glu Asp Asn Lys Lys Leu Ile Ser	65	70	75	80	
gaa act ctg gca acc ttt ggg ggc ctg gat att gtt tgt gct aat gca	85	90	95		288
Glu Thr Leu Ala Thr Phe Gly Gly Leu Asp Ile Val Cys Ala Asn Ala					
gga att gga aag ttc gct ccc acc cat gaa aca ccc ttc gac gta tgg	100	105	110		336
Gly Ile Gly Lys Phe Ala Pro Thr His Glu Thr Pro Phe Asp Val Trp					
aag aag gtg att gct gtg aat ttg aat gga gta ttc tta ctg gat aag	115	120	125		384
Lys Lys Val Ile Ala Val Asn Leu Asn Gly Val Phe Leu Leu Asp Lys					
cta gcc atc aat tac tgg cta gag aaa agc aaa ccc ggc gta att gtc	130	135	140		432
Leu Ala Ile Asn Tyr Trp Leu Glu Lys Ser Lys Pro Gly Val Ile Val					
aac atg gga tca gtc cac tct ttt gta gca gct cct ggc ctt gcg cat	145	150	155	160	480
Asn Met Gly Ser Val His Ser Phe Val Ala Ala Pro Gly Leu Ala His					
tat gga gct gca aaa ggc ggt gtc aaa ctg tta aca caa aca ttg gct	165	170	175		528
Tyr Gly Ala Ala Lys Gly Gly Val Lys Leu Leu Thr Gln Thr Leu Ala					
cta gag tac gca tct cat ggt att aga gta aat tct gtc aat ccg ggg	180	185	190		576
Leu Glu Tyr Ala Ser His Gly Ile Arg Val Asn Ser Val Asn Pro Gly					
tac att tcg act cct ttg ata gat gag gtt ccg aaa gag ccg ttg gat	195	200	205		624
Tyr Ile Ser Thr Pro Leu Ile Asp Glu Val Pro Lys Glu Arg Leu Asp					
aaa ctt gta agc ttg cac cct att ggg aga cta ggt cgt cca gag gaa	210	215	220		672
Lys Leu Val Ser Leu His Pro Ile Gly Arg Leu Gly Arg Pro Glu Glu					
gtt gct gat gca gtc gca ttt ctg tgt tcc cag gag gcc act ttc atc	225	230	235	240	720
Val Ala Asp Ala Val Ala Phe Leu Cys Ser Gln Glu Ala Thr Phe Ile					
aac ggc gtt tct ttg ccg gtt gac ggg ggg tac aca gca cag taa	245	250			765
Asn Gly Val Ser Leu Pro Val Asp Gly Gly Tyr Thr Ala Gln					

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Lys Val Thr Ile Ser Asp Val Ser Gly Glu Lys Lys Tyr His Glu Thr			
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Val Val Ala Leu Lys Ala Gln Asn Leu Asn Thr Asp Asn Leu His Tyr			
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Val Gln Ala Asp Ser Ser Lys Glu Glu Asp Asn Lys Lys Leu Ile Ser			
65	70	75	80
Glu Thr Leu Ala Thr Phe Gly Gly Leu Asp Ile Val Cys Ala Asn Ala			
85	90	95	
Gly Ile Gly Lys Phe Ala Pro Thr His Glu Thr Pro Phe Asp Val Trp			
100	105	110	
Lys Lys Val Ile Ala Val Asn Leu Asn Gly Val Phe Leu Leu Asp Lys			
115	120	125	
Leu Ala Ile Asn Tyr Trp Leu Glu Lys Ser Lys Pro Gly Val Ile Val			
130	135	140	
Asn Met Gly Ser Val His Ser Phe Val Ala Ala Pro Gly Leu Ala His			
145	150	155	160
Tyr Gly Ala Ala Lys Gly Gly Val Lys Leu Leu Thr Gln Thr Leu Ala			
165	170	175	
Leu Glu Tyr Ala Ser His Gly Ile Arg Val Asn Ser Val Asn Pro Gly			
180	185	190	
Tyr Ile Ser Thr Pro Leu Ile Asp Glu Val Pro Lys Glu Arg Leu Asp			
195	200	205	
Lys Leu Val Ser Leu His Pro Ile Gly Arg Leu Gly Arg Pro Glu Glu			
210	215	220	
Val Ala Asp Ala Val Ala Phe Leu Cys Ser Gln Glu Ala Thr Phe Ile			
225	230	235	240
Asn Gly Val Ser Leu Pro Val Asp Gly Gly Tyr Thr Ala Gln			
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 caaaatctca acactgacaa cctccattat gtacaggcag attccagcaa agaagaagat 180  
 aacaagaaat tgatttcgga aactctggca acctttgggg gcctggatat tgtttgc 240  
 aatgcaggaa ttggaaagtt cgctcccacc catgaaacac ccttcgacgt atggaaagaag 300  
 gtgattgctg tgaatttggaa tggagtttgc ttactggata agcttagccat caattactgg 360  
 ctagagaaaa gcaaaccgg cgtaattgtc aacatggat cagtcactc ttttgc 420  
 gctcctggcc ttgcgcattt tggagctgca aaaggcggtg tcaaactgtt aacacaaaca 480  
 ttggctctag agtacgcac tcatggatt agatgaaatt ctgtcaatcc ggggtacatt 540  
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 gatgcacttt tcgagaacac acctgagttac aaaacaatatt atatcattat attagaacag 240  
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 acacctgatt aaaaaatccg gatattaaga atcatgaaac aaaattcaat gttaccctac 360  
 ccattccttc tcggAACCTC ctgtatgactt attaataatgtt aggttgttcc gataaaaaatc 420  
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 aggaggccac tttcatcaac ggcgtttctt tgccgggtga cgggggggtac acagcccgat 180  
 aaattggaca ctttttgctc ttatattatct tccccgcgtt tcaccaatta tccgggtgtac 240  
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 atcagctcgat taaaattatct tggatataataaataagac agaaaaccctgt tggactccta 360  
 gtaggtgtt ctactttcat taaggcagtc acaaaagcaa tggcgaatc aactgatggaa 420  
 aagatagtttta cactggagga gcaggcctac aatggccac ccgcacggat cataggagaa 480  
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27

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42

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34